IN THE SPECIFICATION:

Please insert the enclosed paper copy of the Sequence Listing after the last page of the specification after the Abstract.

Please replace paragraph number **[0030]** with the following paragraph:

[0030] FIGS. 8A and 8B are the multiple nucleotide sequence alignment of human (hCCR12) (SEQ ID NO: 31) or mouse (mcCCR12) (SEQ ID NO: 32) chemokine receptor sequences herein addressed as CCR12 or CCR11.

Please replace paragraph number **[0041]** with the following paragraph:

[0041] Primers to amplify the full-length sequence for mouse CCR12 have been chosen according to the sequence for L-CCR (Accession number: AB009384). The full-length mouse CCR12 coding sequence was amplified from cDNA derived from LPS-stimulated microglia with the following primers: forward, 5'-TATCAAGCAACCTGCCTCAA (SEQ ID NO:[[__]] 1); backward 5'-TGGCATAAAACAATGTGAAGAGA (SEQ ID NO:[[__]] 2).

Please replace paragraph number **[0042]** with the following paragraph:

[0042] Sequence similarity searches using the mouse CCR12 sequence and human databases gave high homology of mouse CCR12 with the human orphan chemokine receptor CRAM-B (Accession number: AF015525). The following primers were designed to get the full-length sequence for the human CCR12. Forward, 5'-CCCAGTGGGCAGTCTGAA (SEQ ID NO:[[__]] 3); backward, 5'-CTTGCATTTGGTGGATGCTA (SEQ ID NO:[[__]] 4).

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Please replace Table 1 and replace it with the following Table 1:

Gene	Primer sequences (5'-3')	PCR product (bp)
CCR1	GTGGTGGGCAATGTCCTAGT (SEQ ID NO:[[__]] 5) TCAGATTGTAGGGGGTCCAG (SEQ ID NO:[[__]] 6)	658
CCR2	GTATCCAAGAGCTTGATGAAGGG (SEQ ID NO:[[__]] 7) GTGTAATGGTGATCATCTTGTTTGGA (SEQ ID NO:[[__]] 8)	532
CCR3	GCACCACCCTGTGAAAAAGT (SEQ ID NO:[[__]] 9) CGAGGACTGCAGGAAAACTC (SEQ ID NO:[[__]] 10)	521
CCR4	AGGCAAGGACCCTGACCTAT (SEQ ID NO:[[__]] 11) GGACTGCGTGTAAGAGGAGC (SEQ ID NO:[[__]] 12)	644
CCR5	ATTCTCCACACCCTGTTTCG (SEQ ID NO:[[__]] 13) TCAGGCTTGTCTTGCTGGAA (SEQ ID NO:[[__]] 14)	350
CCR6	GTGGTGATGACCTTTGCCTT (SEQ ID NO:[[__]] 15) AGGAGGACCATGTTGTGAGG (SEQ ID NO:[[__]] 16)	656
CCR7	AACGGGCTGGTGATACTGAC (SEQ ID NO:[[__]] 17) ATGAAGACTACCACCACGGC (SEQ ID NO:[[__]] 18)	596
CCR8	TTCCTGCCTCGATGGATTAC (SEQ ID NO:[[__]] 19) GCTTCCACCTCAAAGACTGC (SEQ ID NO:[[__]] 20)	591
D6	TCTTCATCACCTGCATGAGC (SEQ ID NO:[[__]] 21) TATGGGAACCACAGCATGAA (SEQ ID NO:[[__]] 22)	400
CCR12	CTGGCGGTGTTTATCTTGGT (SEQ ID NO:[[__]] 23) AACCAGCAGAGGAAAAGCAA (SEQ ID NO:[[__]] 24)	489
GAPDH	CATCCTGCACCACCAACTGCTTAG (SEQ ID NO:[[__]] 25) GCCTGCTTCACCACCTTCTTGATG (SEQ ID NO:[[__]] 26)	346

Please replace Table 5 and replace it with the following Table 5:

Gene		Primer	Annealing
MCP-1	Fw	GTCTCTGTCACGCTTCTGG (SEQ ID NO:[[__]] 27)	56°C
	Rev	GATCTCTCTCTTGAGCTTGG (SEQ ID NO:[[__]] 28)	
CCR2	Fw	GTATCCAAGAGCTTGATGAAGGG (SEQ ID NO:[[__]] 7)	56°C
	Rev	GTGTAATGGTGATCATCTTGTTTGGA (SEQ ID NO:[[__]] 8)	
CCR11	Fw	CTGGCGGTGTTTATCTTGGT (SEQ ID NO:[[__]] 29)	56°C
	Rev	AACCAGCAGAGGAAAAGCAA (SEQ ID NO:[[__]] 30)	
GAPDH	Fw	CATCCTGCACCACCAACTGCTTAG (SEQ ID NO:[[__]] 25)	60°C
	Rev	GCCTGCTTCACCACCTTCTTGATG (SEQ ID NO:[[__]] 26)	

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